

XX Example 1; Fig 1a; 39PP; English.
PS
XX This invention describes novel heparin binding molecules (I). The
CC molecules (I) are useful as heparin antagonist drugs for cardiovascular
CC application and specifically neutralize heparin's conventional
CC anticoagulant properties. (I) are also useful for counteracting actions
CC of heparin locally e.g. in bleeding wounds, vascular anastomoses or
CC leaking prosthetic vascular grafts. (I) is also useful combined in a
CC pharmaceutical composition with insulin as a substitute for protamine
CC for use in treating diabetics. The heparin binding molecules (I)
CC specifically neutralize heparin's conventional anticoagulant properties
CC without causing deleterious hemodynamic side-effects or exacerbation of
CC the proliferative vascular response to injury. (I) are short-duration,
CC intravenous drugs to be used in elective or emergency situations which
CC can safely and specifically neutralize heparin's proliferative response
CC to injury. This sequence represents a heparin-binding peptide described
CC in the method of the invention.
XX Sequence 19 AA;
XX SQ

Key	Location/Qualifiers	
Modified-site	19	/note= "Ala is modified by unidentified R1 group"
XX	EP1232754-A2.	
XX	PD 21-AUG-2002.	
XX	PF 14-FEB-2002; 2002EP-0251027.	
XX	PR 14-FEB-2001; 2001US-268410P.	
XX	(COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.	
XX	Harris RB, Wolz RL, Wolz G;	
XX	WPI; 2002-659478/71.	
XX	Use of cationic helix peptides for treatment of sepsis and for the detection and removal of endotoxins	

AA Disclosure: FIG 1A; 18pp; English.
PPS XX This invention describes a novel use of antibacterial and
CCC immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2,
CCC Tetra-Arg Helix 2 or Tris-Arg Helix 3 for the manufacture of a medicament
CCC for the treatment of sepsis and the detection and removal of endotoxins.
CCC The peptides of the invention are used in method for detection
CCC

endotoxin in a sample comprising contacting the sample with a labelled helix peptide and then detecting the presence of any labelled molecule bound to endotoxin. The peptides can also be used in a method for removing endotoxin in a sample which comprises exposing the sample to a helix peptide, bound to a solid support, then collecting the sample. The endotoxin removal may be *in vivo*, or the peptides may be used to form an affinity trap for endotoxins in e.g. dialysis-type treatments, or for removal of endotoxins from plasma fractionation products. They are also used as model frameworks for endotoxin binding from which new analogues may be designed. This sequence represents the peptide Arg Helix #2 which is used in the construction of Bis-Arg Helix #2, a branched chain peptide described in the method of the invention.

query	Match	Score	Length	DB	Pred.	No.	Mismatches	Indels	Gaps
1	ARARRAARRAARRAFA	100.0%	19	19	19	3.6e-10	0	0	0
1	ARARRAARRAARRAFA	100.0%	19	19	19	3.6e-10	0	0	0
1	ARARRAARRAARRAFA	100.0%	19	19	19	3.6e-10	0	0	0

AB71430	standard; peptide; 16 AA.	
AB71430;		
27-NOV-2002	(first entry)	
Peptide Tris-Arg Helix #3 fragment.		
Sepsis; branched chain peptide; antibacterial; immunosuppressive; endotoxin; helix peptide.		
Synthetic.		
Key-Modified-site	Location/Qualifiers	
	16	
	/note- "Ala is modified by unidentified R1 group"	

21-AUG-2002. 14-FEB-2002; 2002EP-0251027. 14-FEB-2001; 2001US-268410P.

(COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.

Harris RB, Wolz RL, Wolz G; WPI; 2002-659478/71.

Use of cationic helix Peptides for treatment of sepsis and for the detection and removal of endotoxins - Disclosure; Fig 1B; 18pp; English.

This invention describes a novel use of antibacterial and immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, Tetra Arg Helix 2 or Tris Arg Helix 3 for the manufacture of a medicament for the treatment of sepsis and the detection and removal of endotoxins. The peptides of the invention are used in a method for detecting endotoxin in a sample comprising contacting the sample with a labelled helix peptide and then detecting the presence of any labelled molecule bound to endotoxin. The peptides can also be used in a method for removing endotoxin in a sample which comprises exposing the sample to a helix peptide, bound to a solid support, then collecting the sample. The endotoxin removal may be *in vivo* or the endotoxin may be *in vitro* form no

CC affinity trap for endotoxins in e.g. dialysis-type treatments, or for
 CC removal of endotoxins from plasma fractionation products. They are also
 CC used as model frameworks for endotoxin binding from which new analogues
 CC may be designed. This sequence represents the peptide Arg Helix #3 which
 CC is used in the construction of the branched chain peptide Tris-Arg Helix
 CC peptide described in the method of the invention.

XX Sequence 16 AA:
 Query Match 84.2%; Score 16; DB 23; Length 16;
 Best Local Similarity 100.0%; Pred. No. 1.2e-07;
 Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 4 ARRARAARRARA 19
 ||||| 1 ARRARAARRARA 16

Db 1 ARRARAARRARA 16
 Key
 Modified-site 1 /note- "This residue has a side chain
 C(O)-NepsilonN(CH2)3-Tris-ArgHelix#3, where
 the Tris-ArgHelix#3 is represented in AAB1431."
 Modified-site 16 /note- "Acylated residue"
 PT EP1232754-A2.
 PR 21-AUG-2002.
 PF 14-FEB-2002; 2002EP-0251027.
 PR 14-FEB-2001; 2001US-268410P.
 PA (COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.
 PT Harris RB, Wolz RL, Wolz G;
 DR WPI; 2002-659478/71.
 XX Use of cationic helix peptides for treatment of sepsis and for the
 detection and removal of endotoxins
 Disclosure: Fig 2; 18pp; English.

XX This invention describes a novel use of antibacterial and
 CC immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2,
 CC Tetra-Arg Helix 2 or Tris-Arg Helix 3 for the manufacture of a medicament
 CC for the treatment of sepsis and the detection and removal of endotoxins.
 CC The peptides of the invention are used in a method for detecting
 CC endotoxin in a sample comprising contacting the sample with a labelled
 CC helix peptide and then detecting the presence of any labelled molecule
 CC bound to endotoxin. The peptides can also be used in a method for
 CC removing endotoxin in a sample which comprises exposing the sample to a
 CC helix peptide, bound to a solid support, then collecting the sample. The
 CC endotoxin removal may be in vivo, or the peptides may be used to form an
 CC affinity trap for endotoxins in e.g. dialysis-type treatments, or for
 CC removal of endotoxins from plasma fractionation products. They are also

CC used as model frameworks for endotoxin binding from which new analogues
 CC may be designed. This sequence represents the peptide Arg Helix #3 which
 CC is used in the construction of the branched chain peptide Tris-Arg Helix
 CC peptide described in the method of the invention.

XX Sequence 15 AA:
 Query Match 78.9%; Score 15; DB 23; Length 15;
 Best Local Similarity 100.0%; Pred. No. 8.5e-07;
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 5 RRAARAARRARA 19
 ||||| 1 RRAARAARRARA 15
 Db
 RESULT 5
 AAY25078
 ID AAY25078 standard; peptide; 11 AA.
 XX
 AC AAY25078;
 XX 24-AUG-1999 (first entry)
 DE Transduction protein peptide motif 3.
 XX
 KW Anti-pathogen; fusion protein; protein transduction domain; PTD; AET;
 KW cytoxic domain; suppressor; infection; medicament; ddi; ddc; DAT; 3TC;
 KW FTC; DADP; 1592089; CS92; acyclovir; ganciclovir; penciclovir; interferon;
 KW apoptosis; virus; HIV; cytomegalovirus; CMV; herpes simplex virus; HSV-1;
 KW hepatitis virus; Kaposi's sarcoma-associated herpes virus; KSHV;
 KW herpes virus; yellow fever virus; flavivirus; rhinovirus; plasmoidal;
 KW transduction efficiency; cytotoxin.
 XX Unidentified.
 OS
 PN W09929721-A1.
 XX
 PD 17-JUN-1999.
 XX
 PF 10-DEC-1998; 98WO-US26358.
 XX
 PR 20-APR-1998; 9805-0082402.
 PR 10-DEC-1997; 9705-0669012.
 XX
 PA (UNIW) UNIV WASHINGTON.
 XX
 PI Dowdy SF;
 XX
 DR WPI; 1999-394958/33.
 XX
 New anti-pathogen systems, particularly for virus and plasmidium
 PT infections
 XX
 PS Claim 69; Page 37; 123pp; English.
 XX
 This invention describes a novel anti-pathogen system (APS) comprising a
 CC fusion protein constructed from a covalently linked protein transduction
 CC domain (PTD) and a cytotoxic domain. The APS can be used for suppressing
 CC a pathogen infection in a mammal. The method may further comprise
 CC administering a medicament e.g. AET, ddi, ddc, DAT, 3TC, FTC, DADP,
 CC 1592089, CS92, acyclovir, ganciclovir, penciclovir or an interferon. The
 CC APS can also be administered to a mammal in the presence of a pathogen to
 CC induce apoptosis in a predetermined population of cells. The products can
 CC be used for treating mammals suffering from or, susceptible to a viral
 CC infection or a disease associated with a virus, e.g. HIV, cytomegalovirus
 CC (CMV), herpes simplex virus, e.g. type 1 (HSV-1) (HSV-1), Kaposi's virus, type C
 CC (KCV), Kaposi's sarcoma-associated herpes virus (KSHV) or human herpes
 CC virus 8). Yellow fever virus, flavivirus or rhinovirus, or suffering from
 CC or susceptible to plasmidial infection or a disease associated with a
 CC plasmidial infection, e.g. P. falciparum, P. vivax, P. ovale, or
 CC P. malariae. The APS exhibits high transduction efficiency and
 CC specifically kills or injures cells infected by one or more pathogens.

Formation of the cytotoxin is minimized or eliminated in uninfected cells and in infected cells that keep the pathogen inactive. The APS can be specifically tailored to kill or injure cells infected by one or more pathogen strains. This sequence represents a transduction protein motif described in the invention.

Sequence 11 AA:

Query Match	52.6%	Score 10;	DB 20;	Length 11;
Best Local Similarity	100.0%	Pred. No.	0.014;	
Matches	10;	Conservative	0;	Mismatches 0;
Qy	1	ARAARRARA	10	
Db	2	ARAARRARA	11	

RESULT 6

AAB29419
ID AAB29419 standard; Peptide; 11 AA.
XX
AC AAB29419;

XX DT 09-FEB-2001 (first entry)
XX DE Synthetic transduction peptide, SEQ ID NO:6.
XX

XX Protein transduction domain; fusion molecule; therapeutic agent;

XX drug targeting; drug discovery; cell transduction; bioavailability; vaccine; nervous system disorder; Alzheimer's disease; pre-senile dementia; epilepsy; Parkinson's disease; Huntington's disease; pre-senile dementia; meningo; encephalitis; ischaemia; spongiform encephalopathy; dyslexia; age-related memory loss; Lou Gehring's disease; viral infection; HIV; bacterial infection. XX Synthetic.
XX OS WO200062067-A1.
XX PN 19-OCT-2000.
XX PD 28-FEB-2000; 2000060-05097.
XX PR 28-FEB-1999; 9905-0122757.
XX PR 29-AUG-1999; 9905-0151291.
XX PA (UNIW) UNIV WASHINGTON.
XX DR 2000-647439/62.

XX PT Fusion molecules comprising protein transduction domains and therapeutic agents, useful for treating e.g. Alzheimer's and Parkinson's diseases, dementia and epilepsy -
XX PS Claim 36; Page 147; 191pp; English.
XX PT The invention relates to a novel fusion molecule comprising at least one protein transduction domain (PTD) and at least one linked molecule, where the linked molecule has therapeutic or prophylactic activity against a medical condition. The invention also relates to methods of drug discovery in which the test compound is linked to a suitable transducing protein and introduced to a cell; method of killing resistant microorganisms using a suitable fusion molecule; a mammal comprising a covalently linked fusion molecule; and a mammal adapted for experimental use in which at least one transduction molecule has been transduced into essentially all the cells of the mammal. The fusion molecule is used to deliver a therapeutic agent to a mammal, especially a human. The linked molecule may be a vaccine, an anti-infective drug, a cardiovascular drug, an antitumour drug, an analgesic, an antiinflammatory, a diagnostic marker or a drug for the treatment or prevention of a central or peripheral nervous system disorder. The

CC central nervous system (CNS) disorder is especially Alzheimer's disease, Parkinson's disease, Huntington's disease, and also includes pre-senile dementia, epilepsy and seizures, compulsive behaviour, meningitis, (including viral and bacterial meningitis), encephalopathies, ischaemia, CC scrapie (or related spongiform encephalopathies), dyslexia, age-related CC memory loss or Lou Gehring's disease. Fusion molecules can also be CC used to kill virally infected cells, especially those infected with HIV. CC The vaccines are used to treat or prevent bacterial or viral infections. CC The methods are a highly effective means for transducing a molecule CC into an entire mammal or into specific cells tissues, organs and CC systems within it. They also overcome bioavailability problems that CC are associated with many therapeutic agents (e.g., large molecular size, CC hydrophobicity, hydrophilicity, biological resistance), by providing CC efficient transduction of the target cell. The present sequence represents a specifically claimed protein transduction domain.
XX SQ Sequence 11 AA;

RESULT 6
Query Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

RESULT 7

AAY93547
ID AAY93547 standard; Peptide; 11 AA.
XX
AC AAY93547;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;
XX Protein Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;

XX Protein Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;

XX Protein Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;

XX Protein Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;

XX Protein Match 52.6%; Score 10; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ARAARRARA 10
Db 2 ARAARRARA 11

XX Sequence 11 AA;

CC prostate cancer.
 XX Sequence 11 AA;
 SQ Query Match 52.6%; Score 10; DB 21; Length 11;
 Best Local Similarity 100.0%; Pred. No. 0.014;
 Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Oy 1 ARARRARA 10
 |||||||
 Db 2 ARARRARA 11
 RESULT 8
 AAE05278 AAE05278 standard; peptide; 11 AA.
 ID XX
 AAU76085 standard; peptide; 11 AA.
 ID XX
 AAU76085;
 ID XX
 08-MAY-2002 (first entry)
 DT XX
 Peptide transport moiety #4.
 DS XX
 KW Nociceptin; opioid receptor-like 1; ORL1; hypoatraemia;
 KW coronary heart failure; diuretic; thiazide; loop diuretic;
 KW water diuresis; congestive heart failure; liver cirrhosis;
 KW nephrotic syndrome; hypertension; multiple organ failure;
 KW acute renal failure; hypoalaemia; oedema; transport moiety.
 XX
 OS Synthetic.
 OS XX
 WO200198324-A1.
 AC XX
 AAE05278;
 AC XX
 12-SEP-2001 (first entry)
 DT XX
 DE Human immunodeficiency virus (HIV) TAT mutant peptide #5.
 XX
 DNA recombinase domain; protein transduction domain; PTD; mutant;
 KW gene alteration; TAT protein; mutein; Human immunodeficiency virus;
 HIV.
 XX
 OS Human immunodeficiency virus.
 OS Synthetic.
 XX
 WO200149832-A2.
 PN XX
 WO200149832-A2.
 PD XX
 12-JUL-2001.
 XX
 PF 03-JAN-2001; 2001WO-EP000060.
 XX
 PR 07-JAN-2000; 2000EP-0100351.
 PR 10-NOV-2000; 2000EP-0124595.
 PA (ARTE-) ARTEMIS PHARM GMBH.
 XX
 Schwenk F;
 XX
 WPI; 2001-441873/47.
 PR Using site-specific DNA recombinase domain/protein transduction domain fusion proteins for inducing target gene alterations in organisms or cell cultures.
 XX
 PS Claim 5; Page 71; 85PP; English.
 XX
 The present invention relates to use of fusion proteins comprising a site-specific DNA recombinase domain e.g. Cre and a protein transduction domain (PTD) e.g. the Human immunodeficiency virus (HIV) derived TAT Peptide, for preparing an agent for inducing target gene alterations in a living organism or cell culture. The present invention also provides a method for inducing gene alterations in living organisms using the fusion proteins of the invention. The present sequence is a HIV TAT mutant peptide.
 XX
 Sequence 11 AA;
 Query Match 52.6%; Score 10; DB 22; Length 11;
 Best Local Similarity 100.0%; Pred. No. 0.014;
 Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Oy 1 ARARRARA 10
 |||||||
 Db 2 ARARRARA 11
 RESULT 9
 AAU76085

XX PD 10-MAY-2000.
 XX PF 01-OCT-1999; 99EP-0119514.
 XX PR 06-OCT-1998; 98US-0166930.
 XX PA (COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.
 XX PI Harris RB, Sobei M;
 XX DR; 2000-306006/27.
 XX PR New heparin binding molecules, useful for reducing heparin content in a mammal by reducing the anticoagulant effects of heparin.
 XX PS Example 1: Page 8; 39pp; English.
 CC This invention describes novel heparin binding molecules (I). The molecules (I) are useful as heparin antagonist drugs for cardiovascular application and specifically neutralize heparin's counteracting actions of heparin locally e.g. in bleeding wounds, vascular anastomoses or leaking prosthetic vascular grafts. (I) is also useful combined in a pharmaceutical composition with insulin, as a substitute for protamine for use in treating diabetics. The heparin binding molecules (I) specifically neutralize heparin's conventional anticoagulant properties without causing deleterious hemodynamic side-effects or exacerbation of the proliferative vascular response to injury. (I) are short-duration, intravenous drugs to be used in elective or emergency situations which can safely and specifically neutralize heparin's proliferative response to injury. This sequence represents a heparin-binding peptide described in the method of the invention.
 XX Sequence 19 AA;

Query Match 52.6%; Score 10; DB 21; Length 19;
 Best Local Similarity 100.0%; Pred. No. 0.021; Mismatches 0; Indels 0; Gaps 0;

QY 1 ARARRAARA 10
 ||||| | | | |
 Db 10 ARARRAARA 19

RESULT 14
 AAY25077

ID AAY25077 standard; Peptide; 11 AA.

XX AC AAY25077;

XX DT 24-AUG-1999 (first entry)

XX DE Transduction protein peptide motif 2.

XX AC Anti-pathogen; fusion protein; protein transduction domain; PBD; AST;

XX AC cytotoxic domain; suppressor; infection; medicament; ddi; ddc; d4t; 3TC;

XX AC FTc; DAPP; 1592U89; CS2; acyclovir; ganciclovir; penciclovir; interferon;

XX AC apoptosis; virus; HIV; cytomegalovirus; CMV; herpes simplex virus; HSV-1;

XX AC hepatitis virus; Kaposi's sarcoma-associated herpes virus; KSIV;

XX AC herpes virus; yellow fever virus; filovirus; rhinovirus; plasmodial;

XX AC transduction efficiency; cytoxin.

OS Unidentified.

XX PN WO929721-A1.

XX PR 20-APR-1998; 98RS-002402.

XX PR 10-DEC-1997; 97US-0059012.

XX PA (UNIV) UNIV WASHINGTON.

XX PI Dowdy SF;

XX DR WPI; 1999-394958/33.

XX PI Harris RB, Wolz RL, Wolz G;
 XX DR; 2002-659478/71.
 XX PT Use of cationic helix peptides for treatment of sepsis and for the detection and removal of endotoxins -
 XX PR Disclosure; Fig 1A; 18pp; English.
 CC This invention describes a novel use of antibacterial and immunosuppressive peptides designated Arg Helix 2, Bis Arg Helix 2, Tetra Arg Helix 2 or Tris-Arg Helix 3 for the manufacture of a medicament for the treatment of sepsis and the detection and removal of endotoxins. The peptides of the invention are used in a method for detecting endotoxin in a sample comprising contacting the sample with a labelled helix peptide and then detecting the presence of any labelled molecule bound to endotoxin. The peptides can also be used in a method for removing endotoxin in a sample which comprises exposing the sample to a helix peptide, bound to a solid support, then collecting the sample. The endotoxin may be used in vivo, or the peptides may be used to form an affinity trap for endotoxins in e.g. dialysis-type treatments, or for removal of endotoxins from plasma fractionation products. They are also used as model frameworks for endotoxin binding from which new analogues may be designed. This sequence represents the peptide Arg Helix #2 which is used in the construction of Bis-Arg Helix #2, a branched chain peptide described in the method of the invention.
 XX Sequence 19 AA;
 Query Match 52.6%; Score 10; DB 23; Length 19;
 Best Local Similarity 100.0%; Pred. No. 0.021; Mismatches 0; Indels 0; Gaps 0;

QY 1 ARARRAARA 10
 ||||| | | | |
 Db 10 ARARRAARA 19

RESULT 15
 AAY25077

ID AAY25077 standard; Peptide; 11 AA.

XX AC AAY25077;

XX DT 24-AUG-1999 (first entry)

XX DE Transduction protein peptide motif 2.

XX AC Anti-pathogen; fusion protein; protein transduction domain; PBD; AST;

XX AC cytotoxic domain; suppressor; infection; medicament; ddi; ddc; d4t; 3TC;

XX AC FTc; DAPP; 1592U89; CS2; acyclovir; ganciclovir; penciclovir; interferon;

XX AC apoptosis; virus; HIV; cytomegalovirus; CMV; herpes simplex virus; HSV-1;

XX AC hepatitis virus; Kaposi's sarcoma-associated herpes virus; KSIV;

XX AC herpes virus; yellow fever virus; filovirus; rhinovirus; plasmodial;

XX AC transduction efficiency; cytoxin.

OS Unidentified.

XX PN WO929721-A1.

XX PR 17-JUN-1999.

XX PF 10-DEC-1998; 98RS-002402.

XX PR 21-AUG-2002.

XX PF 14-FEB-2002; 2002EP-0251027.

XX PR 14-FEB-2001; 2001US-268410P.

XX PA (COMM-) COMMONWEALTH BIOTECHNOLOGIES INC.

XX New anti-pathogen systems, particularly for virus and plasmodium infections

PT Claim 68; Page 37; 123PP; English.

CC This invention describes a novel anti-pathogen system (APS) comprising a fusion protein constructed from a covalently linked protein transduction domain (PTD) and a cytosolic domain. The APS can be used for suppressing a pathogen infection in a mammal. The method may further comprise administering a medicament e.g. AZT, ddi, dDC, d4T, 3TC, FTC, DADP, APS can also be administered to a mammal in the presence of a pathogen to induce apoptosis in a predetermined population of cells. The products can be used for treating mammals suffering from or susceptible to a viral infection or a disease associated with a virus, e.g. HIV, cytomegalovirus (CMV), herpes simplex virus, e.g. type 1 (HSV-1), hepatitis virus, type C (HCV), Kaposi's sarcoma-associated herpes virus (KSHV) or human herpes virus 8), yellow fever virus, flavivirus or rhinovirus or suffering from or susceptible to plasmodial infection or a disease associated with a plasmodial infection, e.g. P. falciparum, P. ovale, or P. malariae. The APS exhibits high transduction efficiency and specifically kills or injures cells infected by one or more pathogens. Formation of the cytotoxin is minimized or eliminated in uninfected cells and in infected cells that keep the pathogen inactive. The APS can be specifically tailored to kill or injure cells infected by one or more pathogen strains. This sequence represents a transduction protein motif described in the invention.

XX Sequence 11 AA;

Query Match AAB29418 Score 9; DB 20; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.099;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ARARRAAR 9
Db 2 ARARRAAR 10

RESULT 15

AAB29418 ID AAB29418 standard; peptide: 11 AA.
XX DE AC AAB29418;
XX DT 09-FEB-2001 (first entry)

XX Synthetic transduction peptide, SEQ ID NO:5.
XX KW Protein transduction domain; fusion molecule; therapeutic agent; drug targetting; drug discovery; cell transduction; biotavalability; vaccine; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; pre senile dementia; epilepsy; seizure; compulsive behaviour; meningitis; encephalitis; ischaemia; spongiform encephalopathy; dyslexia; age-related memory loss; Lou Gehring's disease; viral infection; HIV; bacterial infection.

XX OS Synthetic.

XX PN WO2000062067-A1.
XX PD 19-OCT-2000.
XX PF 28-FEB-2000; 2000WO-US05097.
XX PR 28-FEB-1999; 99US-0122757.
XX PR 29-AUG-1999; 99US-0151291.
XX PA (UNIV) UNIV WASHINGTON.
XX PI Dowdy SF;
XX

DR WPI; 2000-647439/62.

XX Fusion molecules comprising protein transduction domains and therapeutic agents, useful for treating e.g. Alzheimer's and Parkinson's diseases, dementia and epilepsy -

XX Claim 36; Page 147; 191PP; English.

CC The invention relates to a novel fusion molecule comprising at least one protein transduction domain (PTD) and at least one linked molecule, where the linked molecule has therapeutic or prophylactic activity against a medical condition. The invention also relates to methods of drug discovery in which the test compound is linked to a suitable transducing protein and introduced to a cell; a method of killing resistant microorganisms using a suitable fusion molecule; a mammal comprising a covalently linked fusion molecule; and a mammal adapted for experimental use in which at least one transduction molecule has been transduced into essentially all the cells of the mammal. The fusion molecule is used to deliver a therapeutic agent to a mammal, especially a human. The linked molecule may be a vaccine, an anti-infective drug, a cardiovascular drug, an antitumour drug, an analgesic, an antiinflammatory, a diagnostic marker or a drug for the treatment or prevention of a central or peripheral nervous system disorder. The central nervous system (CNS) disorder is especially Alzheimer's disease, Parkinson's disease, Huntington's disease, and also includes pre-senile dementia, epilepsy and seizures, compulsive behaviour, meningitis (including viral and bacterial meningitis), encephalitis, ischaemia, spongiform encephalopathies), dyslexia, age-related memory loss or Lou Gehring's disease. Fusion molecules can also be used to kill virally infected cells, especially those infected with HIV. The vaccines are used to treat or prevent bacterial or viral infections. The methods are a highly effective means for transducing a molecule into an entire mammal or into specific cells, tissues, organs and systems within it. They also overcome bioavailability problems that are associated with many therapeutic agents (e.g., large molecular size, hydrophobicity, hydrophilicity, biological resistance), by providing efficient transduction of the target cell. The present sequence represents a specifically claimed protein transduction domain.

XX Sequence 11 AA;

Query Match AAB29418 Score 9; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.099;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ARARRAAR 9
Db 2 ARARRAAR 10

Search completed: August 9, 2003, 16:29:05
Job time : 54.7429 secs